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Microbiome. 2018 Dec 4;6:214. doi: [10.1186/s40168-018-0609-y](https://doi.org/10.1186/s40168-018-0609-y)

Correction to: Succession and persistence of microbial communities and antimicrobial resistance genes associated with International Space Station environmental surfaces

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PMCID: PMC6280456 PMID: [30514368](#)

This corrects the article "[Succession and persistence of microbial communities and antimicrobial resistance genes associated with International Space Station environmental surfaces](#)", 204.

Correction to: Microbiome (2018) 6:204

10.1186/s40168-018-0585-2

Following publication of the original article [[1](#)], the authors reported a typographic error in scientific notation in the number of reads, the text should read as:

“Approximately 7.3×10^8 reads associated with microorganisms were generated after high quality trimming from PMA (21 samples) and non-PMA treated (21 samples) samples. All metagenomics reads were normalized across all samples, which yielded 3.1×10^8 in total, and 7.4×10^6 assigned to each sample, without affecting the taxonomic diversity.”

The authors regret these errors and the inconvenience caused.

Reference

1. Singh NK, et al. Succession and persistence of microbial communities and antimicrobial resistance genes associated with International Space Station environmental surfaces. *Microbiome*. 2018;6:204. doi: 10.1186/s40168-018-0585-2. [[DOI](#)] [[PMC free article](#)] [[PubMed](#)] [[Google Scholar](#)]
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